

Heterologous Expression of Terpene Synthase Enzymes in *Methanosarcina*

Grace Van Cott, Darla Brennan, Sean R. Carr, and Nicole R. Buan

University of Nebraska-Lincoln, Department of Biochemistry



Abstract

Methanogens are obligate anaerobes that utilize inexpensive, non-food substrates and make methane as a by-product. Previously in this lab, methanogens have been engineered to direct carbon to isoprene production. Methanogens then could be engineered to make terpenes which are made of several modified isoprene units. Terpenes are molecules of interest to engineer because they are currently synthesized from non-renewable petroleum or are harvested from their endogenous species at low expression levels. Terpenes are part of multi-billion dollar industries like flavoring, fragrance, pharmaceuticals, and have potential in the energy industry.

Methanogens are being investigated for engineering terpenes because they don't exhibit feedback inhibition in their mevalonate (MVA) pathway where terpene precursors come from. A higher percentage of their carbon could be directed to the product of interest than in other model species used for engineering. This project aims to reduce fossil fuel use and increase renewable energy by engineering methanogens to sustainably synthesize terpene compounds.

Research Questions

- Can *Methanosarcina acetivorans* synthesize farnesyl diphosphate, geraniol, linalool, sabinene, pinene, bisabolene, valencene, and humulene from inexpensive non-food feedstocks such as CO₂, methanol, or acetate?
- Can these terpenes be synthesized in appreciable amounts?
- Are these terpenes toxic to *M. acetivorans*?

Terpene Synthases

- Methanogens have a naturally high-flux MVA pathway for isoprenoid lipids compared to model organisms.
- Terpenes are part of billion dollar industries of fragrance and flavoring.
- Terpenes are currently derived from native organisms or petroleum.
- Terpenes have potential in the pharmaceutical, materials, and energy industries:
 - Fuel blending.
 - Sustainable jet fuel with thermal stability and low pressure tolerance.

Figure 1: Chemical structure of terpene compounds of interest for production in *M. acetivorans*.

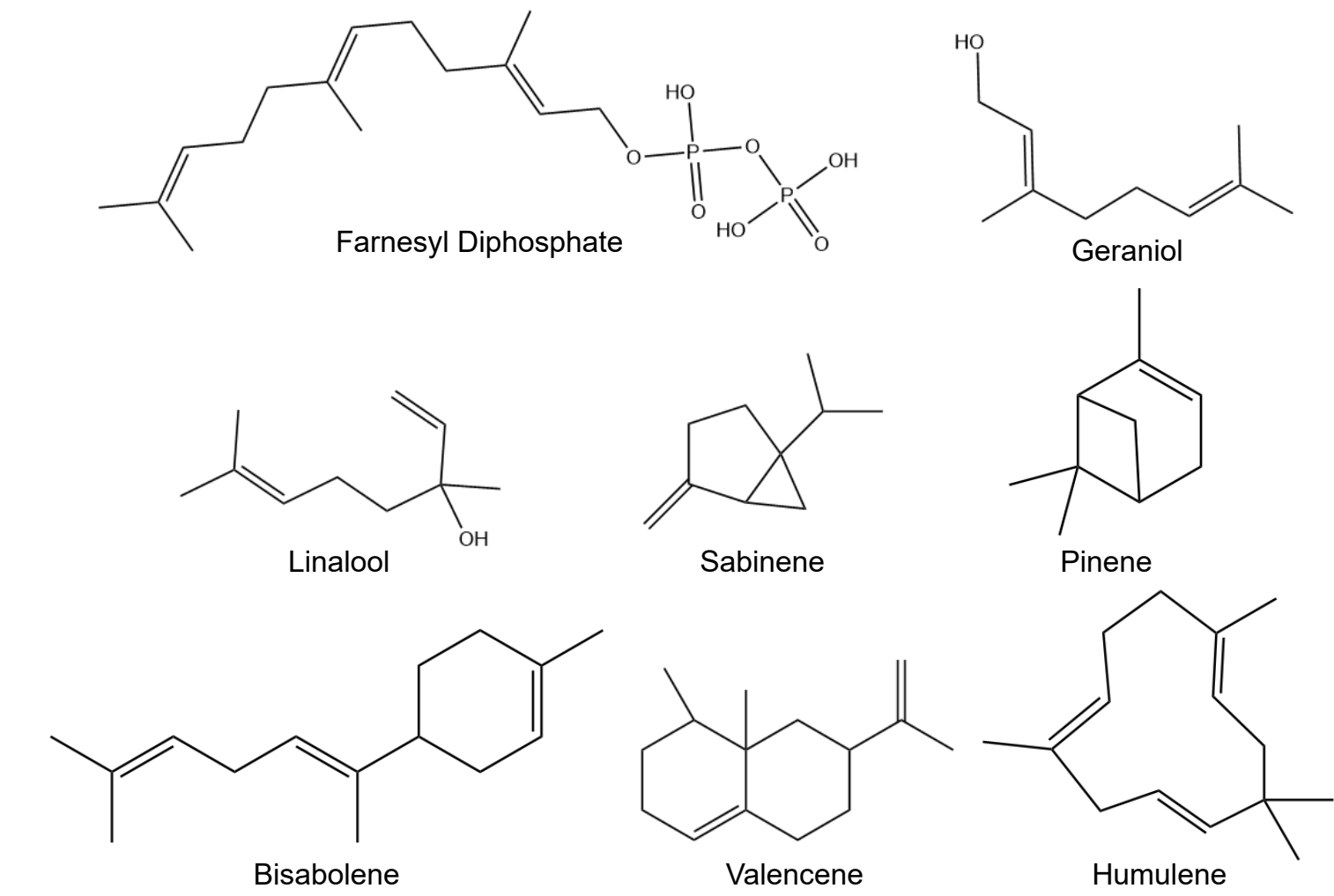
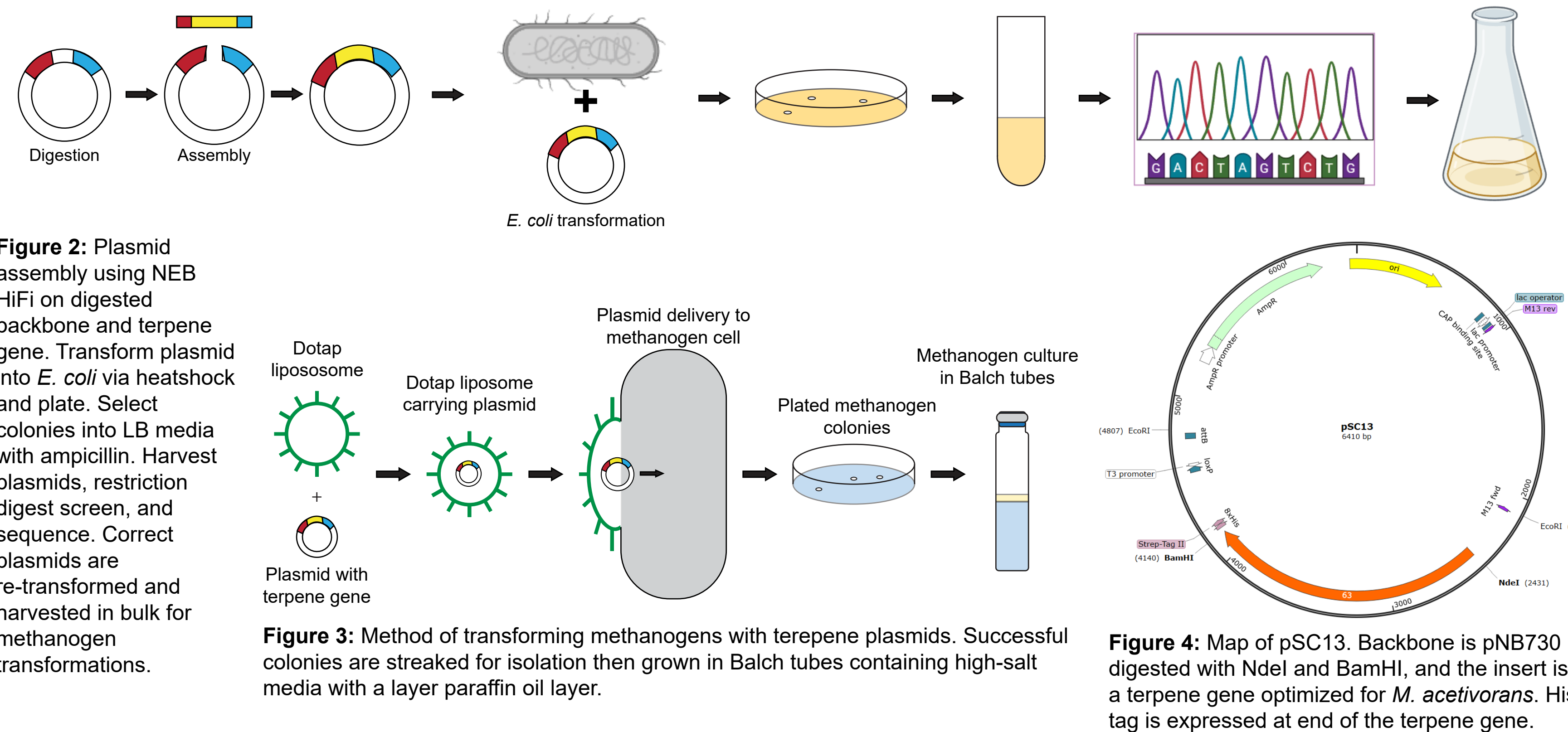


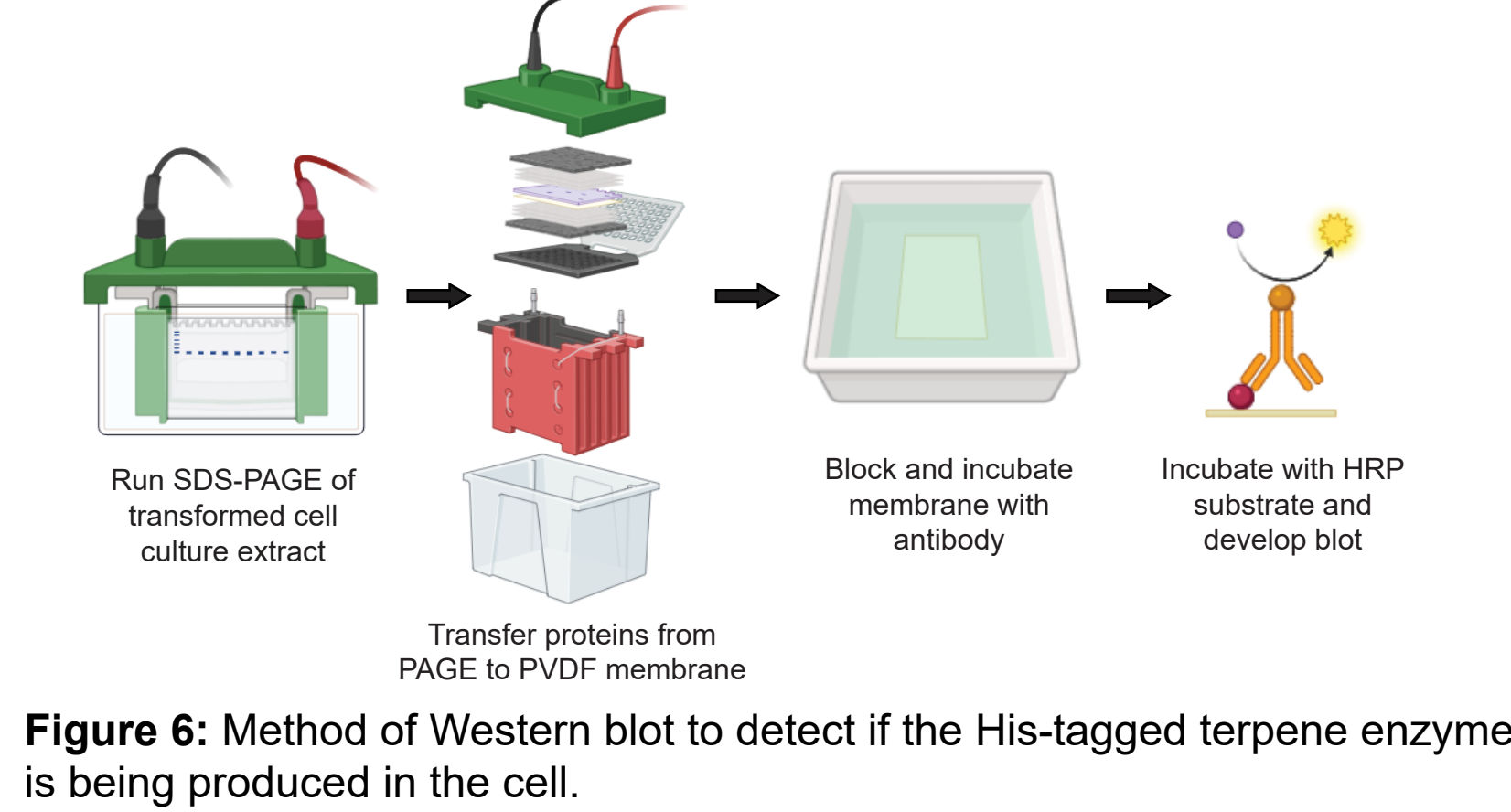
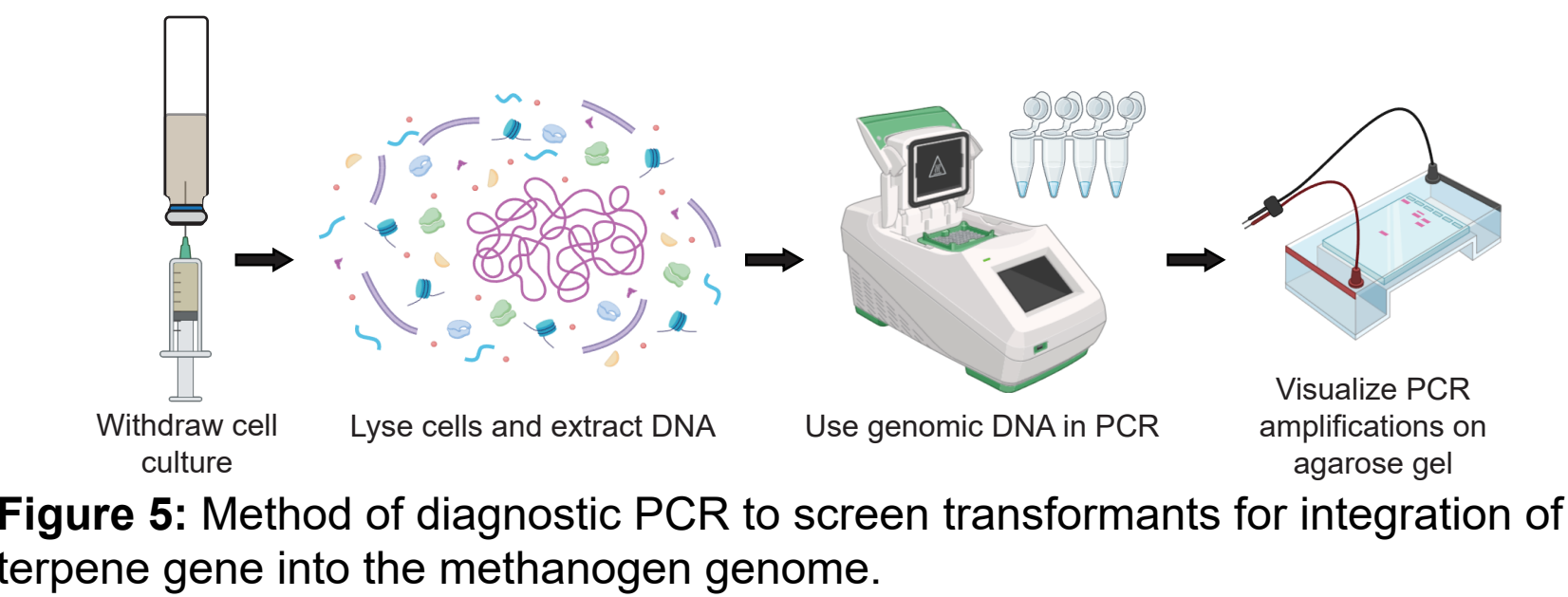
Table 1: Terpene enzyme names, plasmid names, and *M. acetivorans* strain numbers containing the corresponding terpene gene.

Enzyme	Plasmid Name	Strain Number
(2E,6E)-farnesyl diphosphate synthase	pSC12	NB652, NB692
Geraniol synthase	pSC13	NB693, NB695
S-linalool synthase	pSC14	NB697, NB698
(+)-sabinene synthase	pSC17	NB653, NB654
Pinene synthase	pSC18	NB700, NB701
Farnesyl diphosphate synthase	pSC19	NB703, NB704
Alpha-bisabolene synthase	pSC23	NB705, NB706
Valencene synthase	pSC35	NB696
Alpha-humulene synthase	pLK8	NB707, NB708

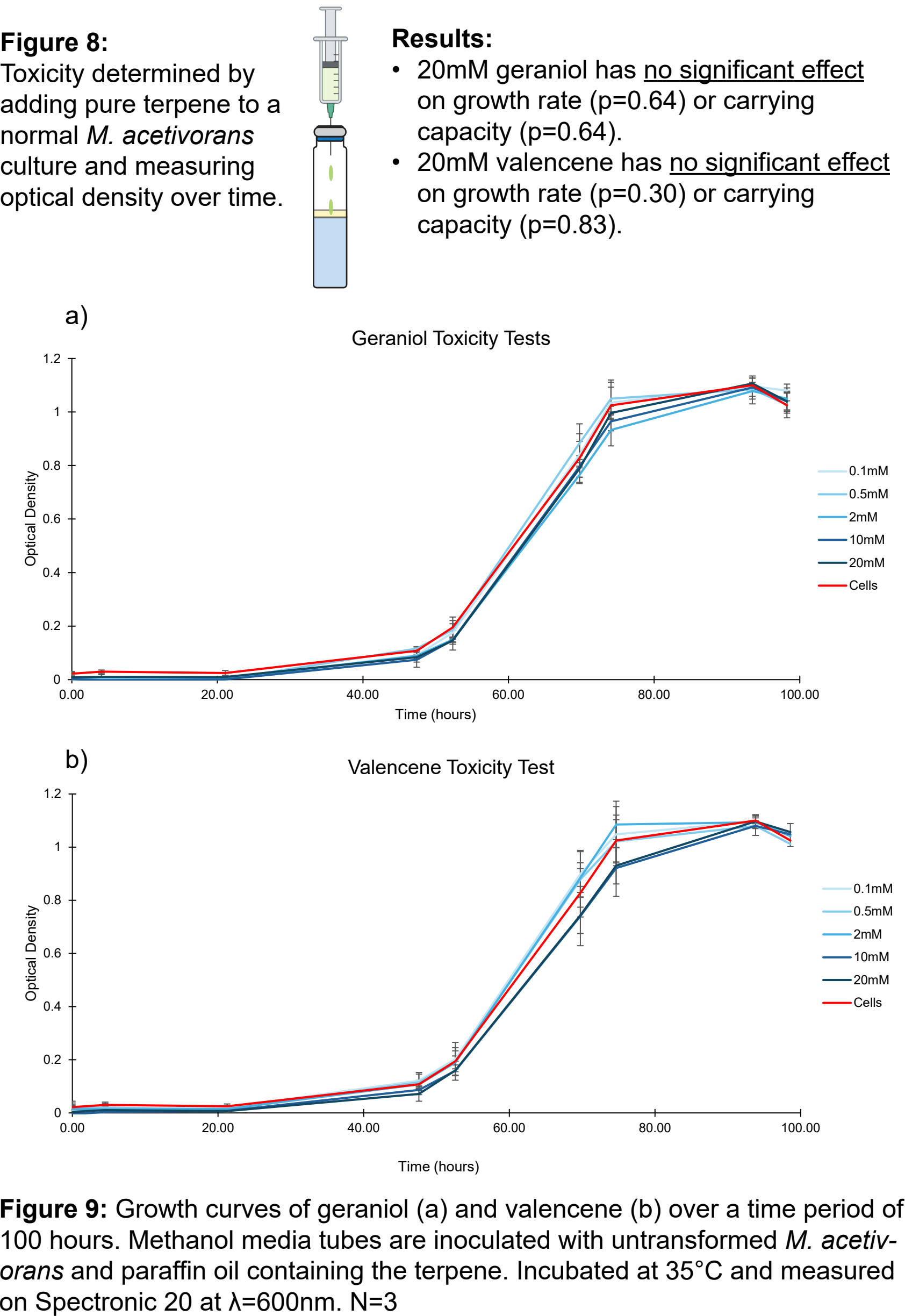
Cloning and Transforming Terpene Expression Plasmids



Strain Validation



Toxicity Tests



Next Steps

- Toxicity tests using farnesyl diphosphate, linalool, sabinene, pinene, bisabolene, and humulene.
- Continue Western blots for farnesyl diphosphate, linalool, sabinene, valencene, pinene, bisabolene, and humulene strains.
- RT-qPCR
- Enzyme Assays
- Gas Chromatography

Figure 10: RT-qPCR procedure to check transcription of the terpene genes.

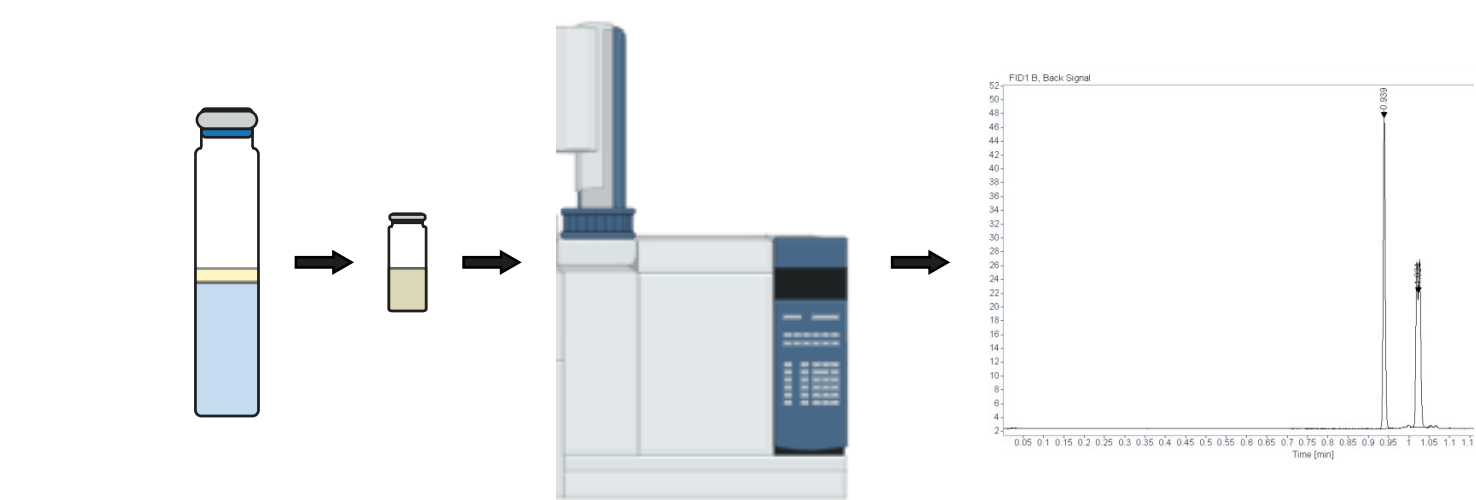


Figure 11: Gas chromatography is performed on the paraffin oil from stationary phase culture. Compare gas chromatogram to the standard terpene in paraffin oil to identify terpene quantities made by methanogens.

Acknowledgements

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References

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